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Sequence Listing was accepted.

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Reviewer: MSpencer

Timestamp: Mon May 07 11:46:59 EDT 2007

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Application No: 10593811

Version No: 1.0

Input Set:**Output Set:**

Started: 2007-04-20 12:04:03.761
Finished: 2007-04-20 12:04:04.552
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 791 ms
Total Warnings: 28
Total Errors: 0
No. of SeqIDs Defined: 44
Actual SeqID Count: 44

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SEQUENCE LISTING

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 Kawamura, Takanori
 Asano, Makoto
 Shitakubo, Daiya
 Shirato, Manabu
 Asada, Makoto

<120> Using non-human animal model, method of measuring transcription
 activity, method of measuring cell number and method of measuring
 tumor volume.

<130> 296514US0PCT

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 <141> 2007-05-07

<150> 10/593,811
 <151> 2006-09-22

<150> PCT/JP05/005257
 <151> 2005-03-23

<150> Japan 2004-084810
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 Gly Ile Ile Pro Val Glu Glu Glu Asn Pro Asp Phe Trp Asn Arg Glu
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 Ala Ala Glu Ala Leu Gly Ala Ala Lys Lys Leu Gln Pro Ala Gln Thr
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 gcc gcc aag aac ctc atc atc ttc ctg ggc gat ggg atg ggg gtg tct 192
 Ala Ala Lys Asn Leu Ile Ile Phe Leu Gly Asp Gly Met Gly Val Ser
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 acg gtg aca gct gcc agg atc cta aaa ggg cag aag aag gac aaa ctg 240
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 65 70 75 80
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 Gly Pro Glu Ile Pro Leu Ala Met Asp Arg Phe Pro Tyr Val Ala Leu
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ctc gtg cat ggc gta cag gaa cag act ttc atc gct cat gta atg gca 1440
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ttc gca gca tgt ttg gag cca tat acc gct tgc gat tta gct cca cca 1488
 Phe Ala Ala Cys Leu Glu Pro Tyr Thr Ala Cys Asp Leu Ala Pro Pro
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Thr Val Thr Ala Ala Arg Ile Leu Lys Gly Gln Lys Lys Asp Lys Leu	65	70	75
Gly Pro Glu Ile Pro Leu Ala Met Asp Arg Phe Pro Tyr Val Ala Leu	85	90	95
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Glu Val Ile Ser Val Met Asn Arg Ala Lys Lys Ala Gly Lys Ser Val	145	150	155
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Tyr Ala His Thr Val Asn Arg Asn Trp Tyr Ser Asp Ala Asp Val Pro	180	185	190
Ala Ser Ala Arg Gln Glu Gly Cys Gln Asp Ile Ala Thr Gln Leu Ile	195	200	205
Ser Asn Met Asp Ile Asp Val Ile Leu Gly Gly Gly Arg Lys Tyr Met	210	215	220
Phe Pro Met Gly Thr Pro Asp Pro Glu Tyr Pro Asp Asp Tyr Ser Gln	225	230	235
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Leu Met Glu Met Thr Glu Ala Ala Leu Arg Leu Leu Ser Arg Asn Pro
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His Glu Ser Arg Ala Tyr Arg Ala Leu Thr Glu Thr Ile Met Phe Asp
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Asp Ala Ile Glu Arg Ala Gly Gln Leu Thr Ser Glu Glu Asp Thr Leu
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Pro Leu Arg Gly Ser Ser Ile Phe Gly Leu Ala Pro Gly Lys Ala Arg
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Ser Pro Glu Tyr Arg Gln Gln Ser Ala Val Pro Leu Asp Glu Glu Thr
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His Ala Gly Glu Asp Val Ala Val Phe Ala Arg Gly Pro Gln Ala His
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Leu Val His Gly Val Gln Glu Gln Thr Phe Ile Ala His Val Met Ala
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 Gly Ile Ile Pro Val Glu Glu Glu Asn Pro Asp Phe Trp Asn Arg Glu
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 Thr Val Thr Ala Ala Arg Ile Leu Lys Gly Gln Lys Lys Asp Lys Leu
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 tcc aag aca tac aat gta gac aaa cat gtg cca gac agt gga gcc aca 336
 Ser Lys Thr Tyr Asn Val Asp Lys His Val Pro Asp Ser Gly Ala Thr
 100 105 110
 gcc acg gcc tac ctg tgc ggg gtc aag ggc aac ttc cag acc att ggc 384
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 115 120 125
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